```
                         GenCore version 5.1.3
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OM protein - nucleic search, using frame_plus_p2n model

Run on:          January 16, 2003, 16:59:22 ; Search time 37.1429 Seconds
                                              (without alignments)
                                              89.447 Million cell updates/sec

Title:           US-09-856-070-18
Perfect score:   24
Sequence:        1 KEELM 5

Scoring table:   BLOSUM62
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                 Ygapop 10.0 , Ygapext  0.5
                 Fgapop  6.0 , Fgapext  7.0
                 Delop   6.0 , Delext   7.0

Searched:        441362 seqs, 153338381 residues

Total number of hits satisfying chosen parameters:      882724

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Command line parameters:
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-Q=/cgn2_1/USPTO_spool/US09856070/runat_14012003_155835_1656/app_query.fasta_1.1592
-DB=Issued_Patents_NA -QFMT=fastap -SUFFIX=rni -MINMATCH=0.1 -LOOPCL=0
-LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
-MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=2000000000
-USER=US09856070_@CGN_1_1_61_@runat_14012003_155835_1656 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database :       Issued_Patents_NA:*
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                 4:  /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
                 5:  /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
                 6:  /cgn2_6/ptodata/2/ina/backfiles1.seq:*

     Pred. No. is the number of results predicted by chance to have a
     score greater than or equal to the score of the result being printed,
     and is derived by analysis of the total score distribution.

                                  SUMMARIES
                    %
Result              Query
  No      Score     Match Length DB  ID                       Description
------------------------------------------------------------------------------
c    1       24     100.0     20  4  US-09-657-481A-28        Sequence 28, Appl
     2       24     100.0     34  4  US-09-291-874-3          Sequence 3, Appli
     3       24     100.0     56  4  US-09-291-874-15         Sequence 15, Appl
     4       24     100.0    539  4  US-09-605-785-310        Sequence 310, App
     5       24     100.0    539  4  US-09-439-313-310        Sequence 310, App
     6       24     100.0    539  4  US-09-352-616A-310       Sequence 310, App
     7       24     100.0    539  4  US-09-232-149A-310       Sequence 310, App
     8       24     100.0    591  2  US-08-332-766A-17        Sequence 17, Appl
     9       24     100.0    728  4  US-08-896-164-53         Sequence 53, Appl
    10       24     100.0    990  1  US-08-410-167A-1         Sequence 1, Appli
    11       24     100.0    993  1  US-08-705-377-1          Sequence 1, Appli
    12       24     100.0    993  1  US-08-705-377-2          Sequence 2, Appli
    13       24     100.0    993  1  US-08-705-377-3          Sequence 3, Appli
    14       24     100.0    993  1  US-08-705-377-4          Sequence 4, Appli
    15       24     100.0    993  1  US-08-705-377-5          Sequence 5, Appli
    16       24     100.0    993  1  US-08-705-377-6          Sequence 6, Appli
    17       24     100.0    993  2  US-09-052-962-1          Sequence 1, Appli
    18       24     100.0    993  2  US-09-052-962-2          Sequence 2, Appli
    19       24     100.0    993  2  US-09-052-962-3          Sequence 3, Appli
    20       24     100.0    993  2  US-09-052-962-4          Sequence 4, Appli
    21       24     100.0    993  2  US-09-052-962-5          Sequence 5, Appli
    22       24     100.0    993  2  US-09-052-962-6          Sequence 6, Appli
    23       24     100.0    993  2  US-09-053-068-1          Sequence 1, Appli
    24       24     100.0    993  2  US-09-053-068-2          Sequence 2, Appli
    25       24     100.0    993  2  US-09-053-068-3          Sequence 3, Appli
    26       24     100.0    993  2  US-09-053-068-4          Sequence 4, Appli
    27       24     100.0    993  2  US-09-053-068-5          Sequence 5, Appli
    28       24     100.0    993  2  US-09-053-068-6          Sequence 6, Appli
    29       24     100.0    993  2  US-08-898-560-2          Sequence 2, Appli
    30       24     100.0    993  4  US-09-101-126-2          Sequence 2, Appli
c   31       24     100.0   1117  4  US-09-177-650-120        Sequence 120, App
    32       24     100.0   1455  1  US-08-446-803-5          Sequence 5, Appli
    33       24     100.0   1455  2  US-08-861-837-5          Sequence 5, Appli
    34       24     100.0   1455  3  US-08-600-656-5          Sequence 5, Appli
    35       24     100.0   1455  4  US-09-170-670-10         Sequence 10, Appl
    36       24     100.0   1455  4  US-09-170-670-15         Sequence 15, Appl
    37       24     100.0   1455  4  US-09-193-068-10         Sequence 10, Appl
    38       24     100.0   1455  4  US-09-193-068-14         Sequence 14, Appl
    39       24     100.0   1455  4  US-09-183-412-10         Sequence 10, Appl
    40       24     100.0   1455  4  US-09-183-412-14         Sequence 14, Appl
    41       24     100.0   1455  4  US-09-354-191A-5         Sequence 5, Appli
    42       24     100.0   1455  4  US-09-290-734-10         Sequence 10, Appl
    43       24     100.0   1455  4  US-09-290-734-15         Sequence 15, Appl
c   44       24     100.0   1689  1  US-07-991-867B-41        Sequence 41, Appl
c   45       24     100.0   1689  2  US-08-544-332-41         Sequence 41, Appl


                                 ALIGNMENTS


RESULT 1
US-09-657-481A-28/c
; Sequence 28, Application US/09657481A
; Patent No. 6258601
; GENERAL INFORMATION:
;  APPLICANT:      Brett P. Monia
;  APPLICANT:  Lex M. Cowsert
;  TITLE OF INVENTION:      ANTISENSE MODULATION OF UBIQUITIN PROTEIN LIGASE WWP1 AND
;  TITLE OF INVENTION:  EXPRESSION
;  FILE REFERENCE      RTS 0087
;  CURRENT APPLICATION NUMBER: US/09/657,481A
;  CURRENT FILING DATE:  2000-09-07
;  NUMBER OF SEQ ID NOS:      93
; SEQ ID NO 28
;   LENGTH: 20
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Antisense Oligonucleotide
US-09-657-481A-28

Alignment Scores:
Pred. No.:              9.02          Length:        20
Score:                  24.00         Matches:       5
Percent Similarity:     100.00%       Conservative:  0
Best Local Similarity:  100.00%       Mismatches:    0
Query Match:            100.00%       Indels:        0
DB:                     4             Gaps:          0

US-09-856-070-18 (1-5) x US-09-657-481A-28 (1-20)

Qy        1 LysGluGluLeuMet 5
              |||||||||||||||
Db       17 AAAGAAGAGTTGATG 3

RESULT 2
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US-09-291-874-3
; Sequence 3, Application US/09291874
; Patent No. 6436694
; GENERAL INFORMATION:
;  APPLICANT: Francis P. Tally
;  APPLICANT: Jianshi Tao
;  APPLICANT: Xiaoyu Shen
;  APPLICANT: Jiansu Zhang
;  TITLE OF INVENTION: Regulable Gene Expression in
;  TITLE OF INVENTION: Gram-Positive Bacteria
;  FILE REFERENCE: CPI98-03p9MA2
;  CURRENT APPLICATION NUMBER: US/09/291,874
;  CURRENT FILING DATE: 1999-04-14
;  EARLIER APPLICATION NUMBER: 60/122,949
;  EARLIER FILING DATE: 1999-03-05
;  EARLIER APPLICATION NUMBER: 09/227,687
;  EARLIER FILING DATE: 1999-01-08
;  EARLIER APPLICATION NUMBER: 60/107,751
;  EARLIER FILING DATE: 1998-11-10
;  EARLIER APPLICATION NUMBER: 60/101,718
;  EARLIER FILING DATE: 1998-09-24
;  EARLIER APPLICATION NUMBER: 60/100,211
;  EARLIER FILING DATE: 1998-09-14
;  EARLIER APPLICATION NUMBER: 60/094,698
;  EARLIER FILING DATE: 1998-07-30
;  EARLIER APPLICATION NUMBER: 60/089,828
;  EARLIER FILING DATE: 1998-06-19
;  EARLIER APPLICATION NUMBER: 60/085,844
;  EARLIER FILING DATE: 1998-05-18
;  EARLIER APPLICATION NUMBER: 60/081,753
;  EARLIER FILING DATE: 1998-04-14
;  EARLIER APPLICATION NUMBER: 60/076,638
;  EARLIER FILING DATE: 1998-03-03
;  EARLIER APPLICATION NUMBER: 60/070,965
;  EARLIER FILING DATE: 1998-01-09
;  NUMBER OF SEQ ID NOS: 28
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 3
;   LENGTH: 34
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Oligonucleotide
US-09-291-874-3

Alignment Scores:
Pred. No.:                16          Length:           34
Score:                    24.00       Matches:          5
Percent Similarity:       100.00%     Conservative:     0
Best Local Similarity:    100.00%     Mismatches:       0
Query Match:              100.00%     Indels:           0
DB:                       4           Gaps:             0

US-09-856-070-18 (1-5) x US-09-291-874-3 (1-34)

Qy        1 LysGluGluLeuMet 5
              |||||||||||||||
Db       12 AAGGAGGAATTAATG 26

RESULT 3
US-09-291-874-15
; Sequence 15, Application US/09291874
; Patent No. 6436694
; GENERAL INFORMATION:
;  APPLICANT: Francis P. Tally
;  APPLICANT: Jianshi Tao
;  APPLICANT: Xiaoyu Shen
;  APPLICANT: Jiansu Zhang
;  TITLE OF INVENTION: Regulable Gene Expression in
;  TITLE OF INVENTION: Gram-Positive Bacteria
;  FILE REFERENCE: CPI98-03p9MA2
;  CURRENT APPLICATION NUMBER: US/09/291,874
;  CURRENT FILING DATE: 1999-04-14
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;  EARLIER APPLICATION NUMBER: 60/122,949
;  EARLIER FILING DATE: 1999-03-05
;  EARLIER APPLICATION NUMBER: 09/227,687
;  EARLIER FILING DATE: 1999-01-08
;  EARLIER APPLICATION NUMBER: 60/107,751
;  EARLIER FILING DATE: 1998-11-10
;  EARLIER APPLICATION NUMBER: 60/101,718
;  EARLIER FILING DATE: 1998-09-24
;  EARLIER APPLICATION NUMBER: 60/100,211
;  EARLIER FILING DATE: 1998-09-14
;  EARLIER APPLICATION NUMBER: 60/094,698
;  EARLIER FILING DATE: 1998-07-30
;  EARLIER APPLICATION NUMBER: 60/089,828
;  EARLIER FILING DATE: 1998-06-19
;  EARLIER APPLICATION NUMBER: 60/085,844
,  EARLIER FILING DATE: 1998-05-18
;  EARLIER APPLICATION NUMBER: 60/081,753
;  EARLIER FILING DATE: 1998-04-14
;  EARLIER APPLICATION NUMBER: 60/076,638
;  EARLIER FILING DATE: 1998-03-03
;  EARLIER APPLICATION NUMBER: 60/070,965
;  EARLIER FILING DATE: 1998-01-09
;  NUMBER OF SEQ ID NOS: 28
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 15
;   LENGTH: 56
;   TYPE: DNA
;   ORGANISM: Artificial Sequence
;   FEATURE:
;   OTHER INFORMATION: Oligonucleotide
US-09-291-874-15

Alignment Scores:
Pred. No.:                27.5        Length:           56
Score:                    24.00       Matches:          5
Percent Similarity:       100.00%     Conservative:     0
Best Local Similarity:    100.00%     Mismatches:       0
Query Match:              100.00%     Indels:           0
DB:                       4           Gaps:             0

US-09-856-070-18 (1-5) x US-09-291-874-15 (1-56)

Qy        1 LysGluGluLeuMet 5
              |||||||||||||||
Db       18 AAGGAGGAATTAATG 32

RESULT 4
US-09-605-785-310
; Sequence 310, Application US/09605785
; Patent No. 6321716
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Mitcham, Jennifer L.
;  APPLICANT:  Harlocker, Susan L.
;  APPLICANT:  Jiang, Yuqui
;  APPLICANT:  Henderson, Robert A.
;  APPLICANT:  Kalos, Michael D.
;  APPLICANT:  Fanger, Gary R.
;  APPLICANT:  Retter, Marc W.
;  APPLICANT:  Stolk, John A.
;  APPLICANT:  Day, Craig H.
;  APPLICANT:  Vedvick, Thomas S.
;  APPLICANT:  Carter, Darrick
;  APPLICANT:  Li, Samuel
;  APPLICANT:  Wang, Aijun
;  APPLICANT:  Skeiky, Yasir A.W.
;  APPLICANT:  Hepler, William
,  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
;  TITLE OF INVENTION:  DIAGNOSIS OF PROSTATE CANCER
;  FILE REFERENCE: 210121.427C16
;  CURRENT APPLICATION NUMBER: US/09/605,785
;  CURRENT FILING DATE:  2000-06-27
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;  NUMBER OF SEQ ID NOS: 835
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 310
;   LENGTH: 539
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-09-605-785-310

Alignment Scores:
Pred. No.:                320            Length:        539
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB:                       4              Gaps:          0

US-09-856-070-18 (1-5) x US-09-605-785-310 (1-539)

Qy       1 LysGluGluLeuMet 5
             ||||||||||||||
Db     464 AAGGAAGAACTTATG 478

RESULT 5
US-09-439-313-310
; Sequence 310, Application US/09439313
; Patent No. 6329505
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Mitcham, Jennifer L.
;  APPLICANT:  Harlocker, Susan Louise
;  APPLICANT:  Jiang Yuqui
;  APPLICANT:  Reed, Steven G.
;  APPLICANT:  Kalos, Michael
;  APPLICANT:  Fanger, Gary
;  APPLICANT:  Retter, Mark
;  APPLICANT:  Solk, John
;  APPLICANT:  Day, Craig
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
;  TITLE OF INVENTION:  DIAGNOSIS OF PROSTATE CANCER
;  FILE REFERENCE: 210121.427C9
;  CURRENT APPLICATION NUMBER: US/09/439,313
;  CURRENT FILING DATE:  1999-11-12
;  NUMBER OF SEQ ID NOS: 575
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 310
;   LENGTH: 539
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-09-439-313-310

Alignment Scores:
Pred. No.:                320            Length:        539
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB:                       4              Gaps:          0

US-09-856-070-18 (1-5) x US-09-439-313-310 (1-539)

Qy       1 LysGluGluLeuMet 5
             ||||||||||||||
Db     464 AAGGAAGAACTTATG 478

RESULT 6
US-09-352-616A-310
; Sequence 310, Application US/09352616A
; Patent No. 6395278
; GENERAL INFORMATION:
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Harlocker, Susan Louise
;  APPLICANT:  Jiang, Yuqui
;  APPLICANT:  Xu, Jiangchun
;  APPLICANT:  Mitcham, Jennifer Lynn
;  TITLE OF INVENTION:  COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
;  TITLE OF INVENTION:  OF PROSTATE CANCER AND METHODS FOR THEIR USE
;  FILE REFERENCE:  210121.427C8
;  CURRENT APPLICATION NUMBER: US/09/352,616A
;  CURRENT FILING DATE:  1999-07-13
;  NUMBER OF SEQ ID NOS:  472
;  SOFTWARE:  FastSEQ for Windows Version 3.0
; SEQ ID NO 310
;   LENGTH: 539
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-09-352-616A-310

Alignment Scores:
Pred. No.:                320            Length:        539
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB:                       4              Gaps:          0

US-09-856-070-18 (1-5) x US-09-352-616A-310 (1-539)

Qy       1 LysGluGluLeuMet 5
             ||||||||||||||
Db     464 AAGGAAGAACTTATG 478

RESULT 7
US-09-232-149A-310
; Sequence 310, Application US/09232149A
; Patent No. 6465611
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Mitcham, Jennifer Lynn
;  TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
;  TITLE OF INVENTION:  CANCER AND METHODS FOR THEIR USE
;  FILE REFERENCE: 210121.427C6
;  CURRENT APPLICATION NUMBER: US/09/232,149A
;  CURRENT FILING DATE:  1999-01-15
;  NUMBER OF SEQ ID NOS: 338
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 310
;   LENGTH: 539
;   TYPE: DNA
;   ORGANISM: Homo sapien
US-09-232-149A-310

Alignment Scores:
Pred. No.:                320            Length:        539
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB:                       4              Gaps:          0

US-09-856-070-18 (1-5) x US-09-232-149A-310 (1-539)

Qy       1 LysGluGluLeuMet 5
             ||||||||||||||
Db     464 AAGGAAGAACTTATG 478

RESULT 8
US-08-332-766A-17
; Sequence 17, Application US/08332766A
; Patent No. 5843647
;  GENERAL INFORMATION:
;    APPLICANT:  JEFFREYS, Alec J.
;    APPLICANT:  ARMOUR, John
;    TITLE OF INVENTION:  SIMPLE TANDEM REPEATS
;    NUMBER OF SEQUENCES:  125
```

```
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  CUSHMAN DARBY & CUSHMAN, L.L.P.
;      STREET:  1100 New York Avenue, N.W.
;      CITY:  Washington
;      STATE:  D. C.
;      COUNTRY:  U.S.A.
;      ZIP:  20005-3918
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/332,766A
;      FILING DATE:  01-NOV-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  GB 9326052.9
;      FILING DATE:  21-DEC-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  BIRD, Donald J.
;      REGISTRATION NUMBER:  25,323
;      REFERENCE/DOCKET NUMBER:  217211/M94/0434/GB
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (202) 861-3000
;      TELEFAX:  (202) 822-0944
;      TELEX:  6714627 CUSH
;  INFORMATION FOR SEQ ID NO:  17:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  591 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-08-332-766A-17

Alignment Scores:
Pred. No.:                354            Length:        591
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB:                       2              Gaps:          0

US-09-856-070-18 (1-5) x US-08-332-766A-17 (1-591)

Qy       1 LysGluGluLeuMet 5
           |||||||||||||||
Db     282 AAACACCAATTAATC 296

RESULT 9
US-08-896-164-53
; Sequence 53, Application US/08896164
; Patent No. 6218521
;  GENERAL INFORMATION:
;    APPLICANT:  OBATA, Yuichi
;    TITLE OF INVENTION:  ISOLATED NUCLEIC ACID MOLECULES ASSOCIATED
;    TITLE OF INVENTION:  WITH GASTRIC CANCER AND METHODS FOR
;    TITLE OF INVENTION:  DIAGNOSING AND TREATING GASTRIC CANCER
;    NUMBER OF SEQUENCES:  87
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Felfe & Lynch
;      STREET:  805 Third Avenue
;      CITY:  New York City
;      STATE:  New York
;      COUNTRY:  USA
;      ZIP:  10022
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette, 3.50 inch, 1.44mb
;      COMPUTER:  IBM PS/2
;      OPERATING SYSTEM:  PC-DOS
;      SOFTWARE:  Wordperfect
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/896,164
;      FILING DATE:  July 17, 1997
;      CLASSIFICATION:  424
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  No. 6218521man D. Hanson
;      REGISTRATION NUMBER:  30,946
;      REFERENCE/DOCKET NUMBER:  LUD 5499 - JEL/NDH/SLH
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 688-9200
;      TELEFAX:  (212) 838-3884
;  INFORMATION FOR SEQ ID NO:  53:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  728 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
US-08-896-164-53

Alignment Scores:
Pred. No.:                444            Length:        728
Score:                    24.00          Matches:       5
Percent Similarity:       100.00%        Conservative:  0
Best Local Similarity:    100.00%        Mismatches:    0
Query Match:              100.00%        Indels:        0
DB.                       4              Gaps:          0

US-09-856-070-18 (1-5) x US-08-896-164-53 (1-728)

Qy       1 LysGluGluLeuMet 5
           |||||||||||||||
Db     232 AAAGAAGAATTAATG 246

RESULT 10
US-08-410-167A-1
; Sequence 1, Application US/08410167A
; Patent No. 5773273
;  GENERAL INFORMATION:
;    APPLICANT:  Tokuzo NISHINO,  Shinichi OHNUMA,  Manabu SUZUKI,
;    APPLICANT:  Chikara OHTO,  Chika ASADA,  Yuka HIGUCHI,  Yoshie TAKEUCHI
;    TITLE OF INVENTION:  Geranylgeranyl-Diphosphate Synthase and DNA
;    TITLE OF INVENTION:  Coding Therefor
;    NUMBER OF SEQUENCES:  4
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kenyon & Kenyon
;      STREET:  One Broadway
;      CITY:  New York
;      STATE:  NY
;      COUNTRY:  US
;      ZIP:  10004
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3+ Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS 6.2
;      SOFTWARE:  WordPerfect 6.1 Windows
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/410,167A
;      FILING DATE:  24-MAR-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 6-53804
;      FILING DATE:  24-MAR-1994
;      APPLICATION NUMBER:  JP 6-315572
;      FILING DATE:  25-NOV-1994
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Edward W. Greason
;      REGISTRATION NUMBER:  18,918
;      REFERENCE/DOCKET NUMBER:
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212)425-7200
;      TELEFAX:  (212)425-5288
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  990 base pairs
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;       TYPE:  nucleic acid
;       STRANDEDNESS:  double
;       TOPOLOGY:  linear
;     MOLECULE TYPE:  DNA (genomic)
;     HYPOTHETICAL:  NO
;     ORIGINAL SOURCE:
;       ORGANISM:  Sulfolobus acidocaldarius
;       STRAIN:  ATCC 33909
;     FEATURE:
;       NAME/KEY:  CDS
;       LOCATION:  1-990
US-08-410-167A-1

Alignment Scores:
Pred. No.:               619          Length:         990
Score:                   24.00        Matches:        5
Percent Similarity:      100.00%      Conservative:   0
Best Local Similarity:   100.00%      Mismatches:     0
Query Match:             100.00%      Indels:         0
DB:                      1            Gaps:           0

US-09-856-070-18 (1-5) x US-08-410-167A-1 (1-990)

Qy       1 LysGluGluLeuMet 5
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Db     811 AAAGAAGAATTAATG 825

RESULT 11
US-08-705-377-1
; Sequence 1, Application US/08705377
; Patent No. 5807725
;  GENERAL INFORMATION:
;     APPLICANT:  OHTO, Chikara,  ASADA, Chika,  OHNUMA, Shinichi,
;     APPLICANT:  NISHINO, Tokuzo,  HIROOKA, Kazutake,  HEMMI, Hisashi
;     TITLE OF INVENTION:  Long Chain Prenyl Diphosphate Synthase
;     NUMBER OF SEQUENCES:  9
;     CORRESPONDENCE ADDRESS:
;       ADDRESSEE:  Kenyon & Kenyon
;       STREET:  1025 Connecticut Avenue, N.W., Suite 600
;       CITY:  Washington
;       STATE:  DC
;       COUNTRY:  USA
;       ZIP:  20036
;     COMPUTER READABLE FORM:
;       MEDIUM TYPE:  3.5 inch Diskette
;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  WordPerfect 6.1 Windows
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/705,377
;       FILING DATE:  29-AUG-1996
;       CLASSIFICATION:  435
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  JP 7247043
;       FILING DATE:  01 SEP 1995
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Toffenetti, Judith L.
;       REGISTRATION NUMBER:  39,048
;       REFERENCE/DOCKET NUMBER:  77670/442
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  202-429-1776
;       TELEFAX:  202-429-0796
;   INFORMATION FOR SEQ ID NO:  1:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  993
;       TYPE:  Nucleic acid
;       STRANDEDNESS:  Double
;       TOPOLOGY:  Linear
;     MOLECULE TYPE:  Genomic DNA
;     ORIGINAL SOURCE:
;       ORGANISM:  Sulfolobus acidocaldarius
US-08-705-377-1
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Alignment Scores:
Pred. No.:               621          Length:         993
Score:                   24.00        Matches:        5
Percent Similarity:      100.00%      Conservative:   0
Best Local Similarity:   100.00%      Mismatches:     0
Query Match:             100.00%      Indels:         0
DB:                      1            Gaps:           0

US-09-856-070-18 (1-5) x US-08-705-377-1 (1-993)

Qy       1 LysGluGluLeuMet 5
             |||||||||||||||
Db     811 AAAGAAGAATTAATG 825

RESULT 12
US-08-705-377-2
; Sequence 2, Application US/08705377
; Patent No. 5807725
;  GENERAL INFORMATION:
;     APPLICANT:  OHTO, Chikara,  ASADA, Chika,  OHNUMA, Shinichi,
;     APPLICANT:  NISHINO, Tokuzo,  HIROOKA, Kazutake,  HEMMI, Hisashi
;     TITLE OF INVENTION:  Long-Chain Prenyl Diphosphate Synthase
;     NUMBER OF SEQUENCES:  9
;     CORRESPONDENCE ADDRESS:
;       ADDRESSEE:  Kenyon & Kenyon
;       STREET:  1025 Connecticut Avenue, N.W., Suite 600
;       CITY:  Washington
;       STATE:  DC
;       COUNTRY:  USA
;       ZIP:  20036
;     COMPUTER READABLE FORM:
;       MEDIUM TYPE:  3.5 inch Diskette
;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  WordPerfect 6.1 Windows
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/705,377
;       FILING DATE:  29-AUG-1996
;       CLASSIFICATION:  435
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  JP 7247043
;       FILING DATE:  01 SEP 1995
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Toffenetti, Judith L.
;       REGISTRATION NUMBER:  39,048
;       REFERENCE/DOCKET NUMBER:  77670/442
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  202-429-1776
;       TELEFAX:  202-429-0796
;   INFORMATION FOR SEQ ID NO:  2:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  993
;       TYPE:  Nucleic acid
;       STRANDEDNESS:  Double
;       TOPOLOGY:  Linear
;     MOLECULE TYPE:  Mutated genomic DNA
US-08-705-377-2

Alignment Scores:
Pred. No.:               621          Length:         993
Score:                   24.00        Matches:        5
Percent Similarity:      100.00%      Conservative:   0
Best Local Similarity:   100.00%      Mismatches:     0
Query Match:             100.00%      Indels:         0
DB:                      1            Gaps:           0

US-09-856-070-18 (1-5) x US-08-705-377-2 (1-993)

Qy       1 LysGluGluLeuMet 5
             |||||||||||||||
Db     811 AAAGAAGAATTAATG 825

RESULT 13
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US-08-705-377-3
; Sequence 3, Application US/08705377
; Patent No. 5807725
;  GENERAL INFORMATION:
;    APPLICANT:  OHTO, Chikara,  ASADA, Chika,  OHNUMA, Shinichi,
;    APPLICANT:  NISHINO, Tokuzo,  HIROOKA, Kazutake,  HEMMI, Hisashi
;    TITLE OF INVENTION:  Long Chain Prenyl Diphosphate Synthase
;    NUMBER OF SEQUENCES:  9
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kenyon & Kenyon
;      STREET:  1025 Connecticut Avenue, N.W., Suite 600
;      CITY:  Washington
;      STATE:  DC
;      COUNTRY:  USA
;      ZIP:  20036
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3.5 inch Diskette
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  WordPerfect 6.1 Windows
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/705,377
;      FILING DATE:  29-AUG-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 7247043
;      FILING DATE:  01 SEP 1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Toffenetti, Judith L.
;      REGISTRATION NUMBER:  39,048
;      REFERENCE/DOCKET NUMBER:  77670/442
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  202-429-1776
;      TELEFAX:  202-429-0796
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  993
;      TYPE:  Nucleic acid
;      STRANDEDNESS:  Double strand
;      TOPOLOGY:  Linear
;    MOLECULE TYPE:  Mutated genomic DNA
US-08-705-377-3

Alignment Scores:
Pred. No.:              621           Length:          993
Score:                  24.00         Matches:         5
Percent Similarity:     100.00%       Conservative:    0
Best Local Similarity:  100.00%       Mismatches:      0
Query Match:            100.00%       Indels:          0
DB:                     1             Gaps:            0

US-09-856-070-18 (1-5) x US-08-705-377-3 (1-993)

Qy       1 LysGluGluLeuMet 5
             |||||||||||||||
Db     811 AAAGAAGAATTAATG 825

RESULT 14
US-08-705-377-4
; Sequence 4, Application US/08705377
; Patent No. 5807725
;  GENERAL INFORMATION:
;    APPLICANT:  OHTO, Chikara,  ASADA, Chika,  OHNUMA, Shinichi,
;    APPLICANT:  NISHINO, Tokuzo,  HIROOKA, Kazutake,  HEMMI, Hisashi
;    TITLE OF INVENTION:  Long-Chain Prenyl Diphosphate Synthase
;    NUMBER OF SEQUENCES:  9
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kenyon & Kenyon
;      STREET:  1025 Connecticut Avenue, N.W., Suite 600
;      CITY:  Washington
;      STATE:  DC
;      COUNTRY:  USA
;      ZIP:  20036
```

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;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3.5 inch Diskette
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  WordPerfect 6.1 Windows
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER.  US/08/705,377
;      FILING DATE:  29-AUG-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 7247043
;      FILING DATE:  01 SEP 1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Toffenetti, Judith L.
;      REGISTRATION NUMBER:  39,048
;      REFERENCE/DOCKET NUMBER:  77670/442
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  202-429-1776
;      TELEFAX:  202-429-0796
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  993
;      TYPE:  Nucleic acid
;      STRANDEDNESS:  Double strand
;      TOPOLOGY:  Linear
;    MOLECULE TYPE:  Mutated genomic DNA
US-08-705-377-4

Alignment Scores:
Pred. No.:              621           Length:          993
Score:                  24.00         Matches:         5
Percent Similarity:     100.00%       Conservative:    0
Best Local Similarity:  100.00%       Mismatches:      0
Query Match:            100.00%       Indels:          0
DB:                     1             Gaps:            0

US-09-856-070-18 (1-5) x US-08-705-377-4 (1-993)

Qy       1 LysGluGluLeuMet 5
             |||||||||||||||
Db     811 AAAGAAGAATTAATG 825

RESULT 15
US-08-705-377-5
; Sequence 5, Application US/08705377
; Patent No. 5807725
;  GENERAL INFORMATION:
;    APPLICANT:  OHTO, Chikara,  ASADA, Chika,  OHNUMA, Shinichi,
;    APPLICANT:  NISHINO, Tokuzo,  HIROOKA, Kazutake,  HEMMI, Hisashi
;    TITLE OF INVENTION:  Long-Chain Prenyl Diphosphate Synthase
;    NUMBER OF SEQUENCES:  9
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Kenyon & Kenyon
;      STREET:  1025 Connecticut Avenue, N.W., Suite 600
;      CITY:  Washington
;      STATE:  DC
;      COUNTRY:  USA
;      ZIP:  20036
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3.5 inch Diskette
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE.  WordPerfect 6.1 Windows
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/705,377
;      FILING DATE:  29-AUG-1996
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  JP 7247043
;      FILING DATE:  01 SEP 1995
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Toffenetti, Judith L.
;      REGISTRATION NUMBER:  39,048
```

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;       REFERENCE/DOCKET NUMBER:  77670/442
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  202-429-1776
;       TELEFAX:  202-429-0796
;   INFORMATION FOR SEQ ID NO:  5:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  993
;       TYPE:  Nucleic acid
;       STRANDEDNESS:  Double strand
;       TOPOLOGY:  Linear
;     MOLECULE TYPE:  Mutated genomic DNA
US-08-705-377-5

Alignment Scores:
Pred. No.:              621            Length:         993
Score:                  24.00          Matches:        5
Percent Similarity:     100.00%        Conservative:   0
Best Local Similarity:  100.00%        Mismatches:     0
Query Match:            100.00%        Indels:         0
DB:                     1              Gaps:           0

US-09-856-070-18 (1-5) x US-08-705-377-5 (1-993)

Qy       1 LysGluGluLeuMet 5
             |||||||||||||||
Db     811 AAAGAAGAATTAATG 825

Search completed: January 16, 2003, 21:41:23
Job time : 19.1429 secs
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